

Table S4. Oligonucleotide sequences used for quantitative PCR analysis of ChIP

Gene (primer location*)	Forward primer (5' to 3')	Reverse primer (5' to 3')
<i>OLFM4</i> (-146 to -128)	TGTGGGCCATGACTCAGATTCC	TATAGTGCCCAGTGGTGATTCC
<i>OLFM4</i> (-1337 to -1190)	CCTAGAGGGGCTAAGTGACATGC	GGAGGTTGCGGCTGTAGTGAG
<i>OLMF4</i> (-11961 to -11783)	CAGTGGCAATGGAAAAGTGA	CAGCTTTGCCCTCTTTACCA
<i>OLFM4</i> (+917 to +1087)	ACCATTTCAGCTCCAACAACAG	CCCTCCAGCCCATTCTCCTTC
<i>HES1</i> [‡] (-105 to +67)	CCTCCCATTGGCTGAAAGTACTG	AGGACCAAGGAGAGAGGTAGACG

*The location of the PCR product is presented as the number of base pairs relative to the transcription start site of the appropriate gene.

[‡]Consensus RBPJκ binding sites are located in the *HES1* promoter at -83 to -76 and -59 to -52 relative to the transcription start site.